

SEQ ID NO: 1

1 LNAGITPVVR EYGLGCSGD LAPLSHCALV LMGEGEA

SEQ ID NO: 2

1 GMLNAGITPV VREYGLGCS GDLAPLSHCA LVLMGEGEAT

SEQ ID NO: 3

1 MASAPQITLG LSGATADDVI AVARHEARIS ISPQVLEELA SVRAHIDALA

51 SADTPVYGIS TGFGALATRH IAPEDRAKLQ RSLIRSHAAG MGEPPERVV

101 RALMFLRAKT LASGRTGVRP VVLETMVGML NAGITPVVRE YGSLGCSGDL

151 APLSHCALVL MGEGEATDAH GDIRPVPELF AEAGLTPVEL AEKEGLALVN

201 GTDGMLGQLI MALADLDELL DIADATAAMS VEAQLGTDQV FRAELHEPLR

251 PHPGQGRSAQ NMFAFLADSP IVASHREGDG RVQDAYS

SEQ ID NO: 4

1 MASAPQITLG LSGATADDVI AVARHEARIS ISPQVLEELA SVRAHIDALA

51 SADTPVYGIS TGFGALATRH IAPEDRAKLQ RSLIRSHAAG MGEPPERVV

101 RALMFLRAKT LASGRTGVRP VVLETMVGML NAGITPVVRE YGSLGCSGDL

151 APLSHCALVL MGEGEATDAH GDIRPVPELF AEAGLTPVEL AEKEGLALVN

201 GTDGMLGQLI MALADLDELL DIADATAAMS VEAQLGTDQV FRAELHEPLR

251 PHPGQGRSAQ NMFAFLADSP IVASHREGDG RVQDAYSLRC SPQVTGAARD

301 TIAHARLVAT RELAAAIDNP VVLPSGEVTS NGNFHGAPVA YVLDFLAIAV

351 ADLGSIAERR TDRMLDPARS RDLPAFLADD PGVDSGMMIA QYTQAGLVAE

401 NKRLA

SEQ ID NO: 5

1	MASAPQITLG	LSGATADDVI	AVARHEARIS	ISPQVLEELA	SVRAHIDALA
51	SADTPVYGIS	TGFGALATRH	IAPEDRAKLQ	RSLIRSHAAG	MGEPVVEREVV
101	RALMFLRAKT	LASGRTGVRP	VVLETMVGML	NAGITPVVRE	YGSLGCSGDL
151	APLSHCALVL	MGEGEATDAH	GDIRPVPELF	AEAGLTPVEL	AEKEGLALVN
201	GTDGMLGQLI	MALADLDELL	DIADATAAMS	VEAQLGTDQV	FRAELHEPLR
251	PHPGQGRSAQ	NMFAFLADSP	IVASHREGDG	RVQDAYSLRC	SPQVTGAARD
301	TIAHARLVAT	RELAAAIDNP	VVLPSGEVTS	NGNFHGAPVA	YVLDFLAIAV
351	ADLGSIAERR	TDRMLDPARS	RDLPAFLADD	PGVDSGMMIA	QYTQAGLVAE
401	NKRLAVPASV	DSIPSSAMQE	DHVS LGWHAA	RKLRTSVANL	RRILAVEMLI
451	AGRALDLRAP	LKPGPATGAV	LEVLRSKVAG	PGQDRFLSAE	LEAAYDLLAN
501	GSVHKALEAH	LPA			

SEQ ID NO: 6

1 XXXXXXXXXXG XSGXTAXDVX AVARHXARXX XSXXXXEXLA XRXXXDALA
51 XXXXPVYGXS TGFGALAXRH IXXEXRAXLQ RXXXRSHAAG MGXXVEREVV
101 RALMFLRXKT XASGXTGVRP XVXXTMXGXL NAGITPVVXE YGSLGCSGDL
151 APLSHCALXL MGEGEATXXX GXXRPXXELX AXXGXXPVEL XEKEGLALXN
201 GTDGM LGXLX MALADLXXLX XXADXTAAXS XEAXLGTDXV XXXELHXXXR
251 PHPGQGSAX NMXXXLAXSX XXXXHXXXXXX RVQDAYSXRC XPQVXGAXRD
301 TXXHAXLVAX RELAXXXDNP VVLPXGXVXS NGNFHGAPVA YVLDFLAIXX
351 ADLGSIXERR TDRXLDXXRS XXLXPFLADD XGVDSGXMIA QYTQAXLVXE
401 XKRLAVPASX DSIPSSAMQE DHVSXGWXAA RKLRTXVXNL XRIXAVEXXX
451 AXRAXXL RAX XXLXPXPAXX AVXXXLRXXX AXGPGXDRFL XXXLXAAXXX
501 XXXGXXXXXAX E

SEQ ID NO: 7

1 ATGGCTTCCG CTCCTCAAAT AACACTTGGC CTAAGTGGCG CAACCGCAGA
51 CGACGTTATC GCCGTTGCCC GCCACGAAGC CCGCATCAGC ATTTCTCCGC
101 AAGTACTTGA GGAAGTGGCT TCCGTCCGAG CACATATCGA TGCACTAGCA
151 TCCGCTGATA CCCC GGTTTA TGGCATTTC A ACCGGCTTTG GCGCGTTGGC
201 AACCCGCCAC ATCGCACCCG AGGATCGCGC CAAGCTGCAG CGCTCCCTCA
251 TCCGTTCCCA CGCTGCTGGC ATGGGTGAAC CGGTGGAGCG CGAAGTGGTC
301 CGCGCATTGA TGTTCCTTGC TGCAAAGACC CTGGCTTCCG GCCGCACGGG
351 CGTTCGCCCC GTTGTCCCTG AGACCATGGT CGGCATGCTC AATGCAGGCA
401 TCACTCCGGT AGTCCGCGAA TACGGTTCAC TGGGCTGCTC CGGTGACTTG
451 GCTCCGCTGT CGCACTGCGC ATTAGTGCTG ATGGGCGAGG GCGAAGCCAC
501 CGATGCCCCAC GGCGACATCC GCCCGGTACC GGAAGTGTTC GCCGAGGCCG
551 GATTGACCCC TGTCGAACTG GCAGAAAAGG AAGGCCTGGC TCTGGTCAAC
601 GGCACCGACG GCATGCTCGG CCAGCTGATC ATGGCATTGG CGGACCTCGA
651 TGAGCTGCTG GACATCGCCG ATGCCACCGC CGCCATGAGC GTTGAAGCCC
701 AGCTGGGCAC CGATCAGGTA TTCCGCGCAG AACTGCACGA ACCACTGCGC
751 CCGCACCCAG GCCAGGGCCG CAGCGCCCAG AACATGTTCG CCTTCCTGGC
801 CGACTCGCCA ATTGTTGCCT CGCATCGCGA GGGAGACGGC CGAGTGCAGG
851 ATGCCTACTC GCTGCGTTGC TCGCCGCAGG TCACCGGCGC CGCCCGCGAC
901 ACCATTGCTC ATGCCCCGCT GGTCGCCACC CGCGAACTGG CTGCGGCCAT

951 TGACAACCCT GTGGTGCTGC CCAGCGGCGA AGTGACTTCC AACGGCAACT

1001 TCCACGGCGC ACCGGTAGCC TACGTGCTGG ACTTCCTTGC CATCGCCGTG

1051 GCCGACCTCG GCTCTATCGC CGAGCGCCGC ACCGACCGCA TGCTCGACCC

1101 AGCCCGCTCC CGCGACCTGC CGGCATTCTT GGCCGACGAT CCGGGTGTGG

1151 ACTCGGGCAT GATGATCGCC CAGTACACCC AGGCCGGCTT GGTGGCAGAA

1201 AACAAAGCGGC TGGCAGTTCC TGCCAGCGTT GACTCCATCC CATCCTCGGC

1251 CATGCAGGAA GACCACGTTT CCCTGGGCTG GCATGCGGCG CGCAAGCTGC

1301 GCACCTCGGT AGCGAACCTC CGCCGCATTC TCGCAGTGGA AATGCTGATT

1351 GCCGGCCGCG CCCTGGACCT GCGGGCCCCA TTGAAGCCTG GTCCAGCGAC

1401 CGGTGCGGTG CTTGAAGTAT TGCGCAGCAA GGTTCAGGC CCCGGCCAGG

1451 ACCGCTTCCT TTCCGCAGAA CTGGAAGCAG CCTATGACCT GCTGGCCAAT

1501 GGCTCGGTGC ATAAGGCCCT CGAAGCTCAC CTGCCTGCAT AA

SEQ ID NO: 8

1 MASAPQITLG LSGATADDVI AVARHEARIS ISPQVLEELA SVRAHIDALA

51 SADTPVYGIS TGFGALATRH IAPEDRAKLQ RSLIRSHAAG MGEPVEREVV

101 RALMFLRAKT LASGRSVRPV VLETMVGMLN AGITPVVREY GSLGCSGDLA

151 PLSHCALVLM GEGEATDAHG DIRPVPELFA EAGLTPVELA EKEGLALVNG

201 TDGMLGQLIM ALADLDELDD IADATAAMSV EAQLGTDQVF RAEHEPLRP

251 HPGQGRSAQN MFAFLADSPI VASHREGDGR VQDAYS

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SEQ ID NO:9

1 MASAPQITLG LSGATADDVI AVARHEARIS ISPQVLEELA SVRAHIDALA

51 SADTPVYGIS TGFGALATRH IAPEDRAKLQ RSLIRSHAAG MGEPVEREVV

101 RALMFLRAKT LASGRSVRPV VLETMVGMLN AGITPVVREY GSLGCSGDLA

151 PLSHCALVLM GEGEATDAHG DIRVPPELFA EAGLTPVELA EKEGLALVNG

201 TDGMLGQLIM ALADLDELDD IADATAAMSV EAQLGTDQVF RAELEPLRP

251 HPGQGRSAQN MFAFLADSPI VASHREGDGR VQDAYSLRCS PQVTGAARDT

301 IAHARLVATR ELAAAIDNPV VLPSGEVTSN GNFHGAPVAY VLDFLAIAVA

351 DLGSIAERRT DRMLDPARSR DLPAFLADDP GVDSGMMIAQ YTQAGLVAEN

401 KRLA

SEQ ID NO:10

1 MASAPQITLG LSGATADDVI AVARHEARIS ISPQVLEELA SVRAHIDALA

51 SADTPVYGIS TGFGALATRH IAPEDRAKLQ RSLIRSHAAG MGEPPEREVV

101 RALMFLRAKT LASGRSVRPV VLETMVGMLN AGITPVVREY GSLGCSGDLA

151 PLSHCALVLM GEGEATDAHG DIRVPPELFA EAGLTPVELA EKEGLALVNG

201 TDGMLGQLIM ALADLDELDD IADATAAMSV EAQLGTDQVF RAEHEPLRP

251 HPGQGRSAQN MFAFLADSPI VASHREGDGR VQDAYSLRCS PQVTGAARDT

301 IAHARLVATR ELAAAIDNPV VLPSGEVTSN GNFHGAPVAY VLDFLAIAVA

351 DLGSIAERRT DRMLDPARSR DLPAFLADDP GVDSGMMIAQ YTQAGLVAEN

401 KRLAVPAVDS IPSSAMQEDH VSLGWHAARK LPTSVANLRR ILAVEMLIAG

451 RALDLRAPLK PGPATGAVLE VLRKVVAGPG QDRFLSAELE AAYDLLANGS

501 VHKALEAHL P E

SEQ ID NO: 11

1 XXXXXXXXXXG XSGXTAXDVX AVARHXARXX XSXXXXEXLA XXRXXDXALA

51 XXXXPVYGXS TGFGALAXRH IXXEXRAXLQ RXXXRSHAAG MGXXVEREVV

101 RALMFLRXKT XASGXX-VRP XVXXTMX**G**XL NAGITPVVXE YGSLGCSGDL

151 APLSHCALVL MGEGEAT**T**XXX GXXRPXXELX AXXGXXPVEL XEKEGLALXN

201 GTDGMLGXLX MALADLXXLX XXADXTAAXS XEAXLGTDXV XXXELHXXXR

251 PHPGQGSAX NMXXXLAX-S XXXXXHXXXX X---RVQDAY SXRCXPQVXG

301 AXRDTXXHAX LVAXRELAXX XDNPVVLXPX -XVXSNGNFH GAPVAYVLDF

351 LAIXXADLGS IXERRTDRXL DXXRSXXLPX FLADDXGVDS GXMIAQYTQA

401 XLVXEXKRLA VPA-XDSIPS SAMQEDHVSX GWXAARKLXT XVXNLXRIXA

451 VEXXXAXRAX XLRAX-XXXX PAXXAVXXXL RXXXA-GPGQ DRFLXXXLXA

501 AXXXXXXXX-- -X-XXXXXEX XXXX

SEQ ID NO: 12

1 ATGGCTTCCG CTCCTCAAAT AACACTTGGC CTAAGTGGCG CAACCGCAGA

51 CGACGTTATC GCCGTTGCCC GCCACGAAGC CCGCATCAGC ATTTCTCCGC

101 AAGTACTTGA GGAAGTGGCT TCCGTCCGAG CACATATCGA TGCACTAGCA

151 TCCGCTGATA CCCCGGTTTA TGGCATTTC A CCGGCTTTG GCGCGTTGGC

201 AACCCGCCAC ATCGCACCCG AGGATCGCGC CAAGCTGCAG CGCTCCCTCA

251 TCCGTTCCCA CGCTGCTGGC ATGGGTGAAC CGGTGGAGCG CGAAGTGGTC

301 CGCGCATTGA TGTTCTTGCG TGCAAAGACC CTGGCTTCCG GCCGCAGCGT

351 TCGCCCGGTT GTCCTTGAGA CCATGGTCGG CATGCTCAAT GCAGGCATCA

401 CTCCGGTAGT CCGCGAATAC GGTTCACTGG GCTGCTCCGG TGACTTGGCT

451 CCGCTGTGCG ACTGCGCATT AGTGCTGATG GGCGAGGGCG AAGCCACCGA

501 TGCCCACGGC GACATCCGCC CGGTACCGGA ACTGTTCGCC GAGGCCGGAT

551 TGACCCCTGT CGAACTGGCA GAAAAGGAAG GCCTGGCTCT GGTCAACGGC

601 ACCGACGGCA TGCTCGGCCA GCTGATCATG GCATTGGCGG ACCTCGATGA

651 GCTGCTGGAC ATCGCCGATG CCACCGCCGC CATGAGCGTT

GAAGCCCAGC

701 TGGGCACCGA TCAGGTATTC CGCGCAGAAC TGCACGAACC ACTGCGCCCG
751 CACCCAGGCC AGGGCCGCAG CGCCAGAAC ATGTTGCCT TCCTGGCCGA
801 CTCGCCAATT GTTGCTCGC ATCGCGAGGG AGACGGCCGA GTGCAGGATG
851 CCTACTCGCT GCGTTGCTCG CCGCAGGTCA CCGGCGCCGC CCGCGACACC
901 ATTGCTCATG CCCGCTGGT CGCCACCCGC GAACTGGCTG CGGCCATTGA
951 CAACCCTGTG GTGCTGCCCA GCGGCGAAGT GACTTCCAAC GGCAACTTCC
1001 ACGGCGCACC GGTAGCCTAC GTGCTGGACT TCCTTGCCAT CGCCGTGGCC
1051 GACCTCGGCT CTATCGCCGA GCGCCGCACC GACCGCATGC TCGACCCAGC
1101 CCGCTCCCGC GACCTGCCGG CATTCCTGGC CGACGATCCG GGTGTGGACT
1151 CGGGCATGAT GATCGCCCAG TACACTCAGG CCGGCTTGGT GGCAGAAAAC
1201 AAGCGGCTGG CAGTTCCTGC AGTTGACTCC ATCCCATCCT CGGCCATGCA
1251 GGAAGACCAC GTTTCCTGG GCTGGCATGC GCGCGCAAG CTGCCGACCT
1301 CGGTAGCGAA CCTCCGCCGC ATTCTCGCAG TGAAATGCT GATTGCCGGC

1351 CGCGCCCTGG ACCTGCGGGC CCCATTGAAG CCTGGTCCAG

CGACCGGTGC

1401 GGTGCTTGAA GTATTGCGCA GCAAGGTTGC AGGCCCCGGC CAGGACCGCT

1451 TCCTTTCCGC AGAACTGGAA GCAGCCTATG ACCTGCTGGC CAATGGCTCG

1501 GTGCATAAGG CCCTCGAAGC TCACCTGCCT GAATAA

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